SEQUENCE LISTING

<110> Hill, Ronald J. Hannan, Garry N.

TECH CENTER 1600/2900

<120> Genetic Sequences Encoding Steroid and Juvenile Hormone Receptor Polypeptides and Insecticidal Modalities Therefor

<130> 53-99

<140> 09/346,470

<141> 1999-07-01

<150> WO 99/00033

<151> 1999-01-15

<150> AU PP1356/98

<151> 1998-01-15 .

<160> 20

<170> PatentIn Ver. 2.0

<210> 1

<211> 2274

<212> DNA

<213> Lucilia cuprina

<220>

<221> CDS

<222> (1)..(2271)

<400> 1

atg atg aaa cga cgt tgg tct aat aat ggc ggt ttt gcc gct tta aaa 48 Met Met Lys Arg Arg Trp Ser Asn Asn Gly Gly Phe Ala Ala Leu Lys 10

atg tta gaa gaa tcc tcc tca gaa gta acc tcc tcc tca aat ggt ctg 96 Met Leu Glu Glu Ser Ser Ser Glu Val Thr Ser Ser Ser Asn Gly Leu 20

gtc ttg tca tcg gat ata aat atg tca cct tcc tcg ttg gat tca ccc 144 Val Leu Ser Ser Asp Ile Asn Met Ser Pro Ser Ser Leu Asp Ser Pro 35 40 45

gtt tat ggc gat cag gaa atg tgg ctg tgt aac gat tca gct tca tat 192 Val Tyr Gly Asp Gln Glu Met Trp Leu Cys Asn Asp Ser Ala Ser Tyr

Page 1 of 27

			cat His 70											240
			caa Gln											288
			tcc Ser											336
			aca Thr											384
 		 	ggt Gly			_		~ ~	_					432
			ggt Gly 150										_	480
			cac His				_			_		_	-	528
			ggc											576
			aat Asn					_	-		_		_	624
			gcc Ala						_			_		672
		-	atc Ile 230		-									720
	_		agt Ser	_	_	_					-		-	768
_			tca Ser			_	_	-				_	_	816

			tca Ser								864
			ccc Pro								912
			ggt Gly 310								960
			cga Arg								1008
		His	gcc Ala								1056
			ctg Leu								1104
			gaa Glu								1152
			aag Lys 390		_	-		_	-	_	1200
			ata Ile								1248
			ctg Leu								1296
			cct Pro								1344
			caa Gln								1392
			agt Ser 470			_	_			_	1440

	ttt Phe									tta Leu	1488
	gaa Glu										1536
	caa Gln 515										1584
	atg Met										1632
	aat Asn		Ser				_	_		_	1680
	aat Asn										1728
	gtg Val				-		-				1776
	gat Asp 595										1824
	tac Tyr				_				_		1872
	gat Asp										1920
	gaa Glu										1968
-	aaa Lys	_		-	_			-			2016
-	gta Val 675		_			 _			_	_	2064

acc Thr	cag Gln 690	Ala	gaa Glu	aag Lys	gcc Ala	gcc Ala 695	cag Gln	gaa Glu	gct Ala	cag Gln	gca Ala 700	aca Thr	aca Thr	tcg Ser	gcc Ala	2112
att Ile 705	tca Ser	gca Ala	gcc Ala	gcc Ala	acc Thr 710	tca Ser	tct Ser	tcc Ser	tcc Ser	ata Ile 715	aat Asn	acc Thr	tcg Ser	atg Met	gca Ala 720	2160
aca Thr	tca Ser	tcc Ser	tca Ser	tca Ser 725	tcg Ser	tta Leu	tcg Ser	cca Pro	tcg Ser 730	gcg Ala	gcc Ala	tca Ser	aca Thr	ccc Pro 735	aat Asn	2208
ggt Gly	ggt Gly	gcc Ala	gtc Val 740	gat Asp	tat Tyr	gtt Val	ggc Gly	acc Thr 745	gat Asp	atg Met	agt Ser	atg Met	agt Ser 750	tta Leu	gta Val	2256
		gat Asp 755		-	tag											2274
<212	L> 7! 2> PI		ia cı	ıpri:	na											
<400)> 2															
		Lys	Arg	Arg 5	Trp	Ser	Asn	Asn	Gly 10	Gly	Phe	Ala	Ala	Leu 15	Lys	
Met 1	Met			5	Trp Ser				10	_				15	-	
Met 1 Met	Met Leu	Glu	Glu 20	5 Ser		Ser	Glu	Val 25	10 Thr	Ser	Ser	Ser	Asn 30	15 Gly	Leu	
Met Met Val	Met Leu Leu	Glu Ser 35	Glu 20 Ser	5 Ser Asp	Ser	Ser Asn	Glu Met 40	Val 25 Ser	10 Thr Pro	Ser	Ser Ser	Ser Leu 45	Asn 30 Asp	15 Gly Ser	Leu Pro	
Met Met Val	Met Leu Leu Tyr 50	Glu Ser 35 Gly	Glu 20 Ser Asp	5 Ser Asp Gln	Ser Ile	Ser Asn Met 55	Glu Met 40 Trp	Val 25 Ser Leu	10 Thr Pro Cys	Ser Ser Asn	Ser Ser Asp	Ser Leu 45 Ser	Asn 30 Asp Ala	15 Gly Ser	Leu Pro Tyr	
Met 1 Met Val Val Asn 65	Met Leu Leu Tyr 50 Asn	Glu Ser 35 Gly Ser	Glu 20 Ser Asp	Ser Asp Gln	Ser Ile Glu His	Ser Asn Met 55 Ser	Glu Met 40 Trp Val	Val 25 Ser Leu Ile	10 Thr Pro Cys	Ser Ser Asn Ser 75	Ser Ser Asp 60 Leu	Ser Leu 45 Ser Gln	Asn 30 Asp Ala Gly	15 Gly Ser Ser	Leu Pro Tyr Thr 80	
Met 1 Met Val Val Asn 65 Ser	Met Leu Leu Tyr 50 Asn	Glu Ser 35 Gly Ser Leu	Glu 20 Ser Asp His	Ser Asp Gln Gln Ala 85	Ser Ile Glu His 70	Ser Asn Met 55 Ser	Glu Met 40 Trp Val	Val 25 Ser Leu Ile	Thr Pro Cys Thr	Ser Ser Asn Ser 75	Ser Ser Asp 60 Leu Leu	Ser Leu 45 Ser Gln Ser	Asn 30 Asp Ala Gly	15 Gly Ser Ser Cys Leu 95	Leu Pro Tyr Thr 80 Pro	

Gly	Gly		Gly	Gly	Gly	Gly 135		Val	Pro	Gly	Met 140	Thr	Ser	Leu	Asn
Gly 145		Gly	Gly	Gly	Gly 150	Gly	Ser	Gln	Val	Asn 155	Asn	His	Asn	His	Ser 160
His	Asn	His	Leu	His 165	His	Asn	Ser	Asn	Ser 170	Asn	His	Ser	Asn	Ser 175	Ser
Ser	His	His	Thr 180	Asn	Gly	His	Met	Gly 185	Ile	Gly	Gly	Gly	Gly 190	Gly	Gly
Leu	Ser	Val 195	Asn	Ile	Asn	Gly	Pro 200	Asn	Ile	Val	Ser	Asn 205	Ala	Gln	Gln
Leu	Asn 210	Ser	Leu	Gln	Ala	Ser 215	Gln	Asn	Gly	Gln	Val 220	Ile	His	Ala	Asn
Ile 225	Gly	Ile	His	Ser	Ile 230	Ile	Ser	Asn	Gly	Leu 235	Asn	His	His	His	His 240
His	His	Met	Asn	Asn 245	Ser	Ser	Met	Met	His 250	His	Thr	Pro	Arg	Ser 255	Glu
Ser	Ala	Asn	Ser 260	Ile	Ser	Ser	Gly	Arg 265	Asp	Asp	Leu	Ser	Pro 270	Ser	Ser
Ser	Leu	Asn 275	Gly	Phe	Ser	Thr	Ser 280	Asp	Ala	Ser	Asp	Val 285	Lys	Lys	Ile
Lys	Lys 290	Gly	Pro	Ala	Pro	Arg 295	Leu	Gln	Glu	Glu	Leu 300	Cys	Leu	Val	Cys
Gly 305	Asp	Arg	Ala	Ser	Gly 310	Tyr	His	Tyr	Asn	Ala 315	Leu	Thr	Cys	Glu	Gly 320
Cys	Lys	Gly	Phe	Phe 325	Arg	Arg	Ser	Val	Thr 330	Lys	Asn	Ala	Val	Tyr 335	Cys
Cys	Lys	Phe	Gly 340	His	Ala	Cys	Glu	Met 345	Asp	Met	Tyr	Met	Arg 350	Arg	Lys
Cys	Gln	Glu 355	Cys	Arg	Leu	Lys	Lys 360	Cys	Leu	Ala	Val	Gly 365	Met	Arg	Pro
Glu	Cys 370	Val	Val	Pro	Glu	Asn 375	Gln	Cys	Ala		Lys 380	Arg	Arg	Glu	Lys
Lys 385	Ala	Gln	Lys	Glu	Lys 390	Asp	Lys	Ile	Gln	Thr 395	Ser	Val	Cys	Ala	Thr 400

Glu	Ile	Lys	Lys	Glu 405	Ile	Leu	Asp	Leu	Met 410	Thr	Cys	Glu	Pro	Pro 415	Ser
His	Pro	Thr	Cys 420	Pro	Leu	Leu	Pro	Glu 425	Asp	Ile	Leu	Ala	Lys 430	Cys	Gln
Ala	Arg	Asn 435	Ile	Pro	Pro	Leu	Ser 440	Tyr	Asn	Gln	Leu	Ala 445	Val	Ile	Tyr
Lys	Leu 450	Ile	Trp	Tyr	Gln	Asp 455	Gly	Tyr	Glu	Gln	Pro 460	Ser	Glu	Glu	Asp
Leu 465	Lys	Arg	Ile	Met	Ser 470	Ser	Pro	Asp	Glu	Asn 475	Glu	Ser	Gln	His	Asp 480
Ala	Ser	Phe	Arg	His 485	Ile	Thr	Glu	Ile	Thr 490	Ile	Leu	Thr	Val	Gln 495	Leu
Ile	Val	Glu	Phe 500	Ala	Lys	Gly	Leu	Pro 505	Ala	Phe	Thr	Lys	Ile 510	Pro	Gln
Glu	Asp	Gln 515	Ile	Thr	Leu	Leu	Lys 520	Ala	Cys	Ser	Ser	Glu 525	Val	Met	Met
Leu	Arg 530	Met	Ala	Arg	Arg	Tyr 535	Asp	His	Asn	Ser	Asp 540	Ser	Ile	Phe	Phe
Ala 545	Asn	Asn	Arg	Ser	Tyr 550	Thr	Arg	Asp	Ser	Tyr 555	Lys	Met	Ala	Gly	Met 560
Ala	Asp	Asn	Ile	Glu 565	Asp	Leu	Leu	His	Phe 570	Cys	Arg	Gln	Met	Tyr 575	Ser
Met	Lys	Val	Asp 580	Asn	Val	Glu	Tyr	Ala 585	Leu	Leu	Thr	Ala	Ile 590	Val	Ile
Phe	Ser	Asp 595	Arg	Pro	Gly	Leu	Glu 600	Glu	Ala	Glu	Leu	Val 605	Glu	Ala	Ile
Gln	Ser 610	Tyr	Tyr	Ile	Asp	Thr 615	Leu	Arg	Ile	Tyr	Ile 620	Leu	Asn	Arg	His
Cys 625	Gly	Asp	Pro	Met	Ser 630	Leu	Val	Phe	Phe	Ala 635	Lys	Leu	Leu	Ser	Ile 640
Leu	Thr	Glu	Leu	Arg 645	Thr	Leu	Gly	Asn	Gln 650	Asn	Ala	Glu	Met	Cys 655	Phe
Ser	Leu	Lys	Leu 660	Lys	Asn	Arg	Lys	Leu 665	Pro	Lys	Phe	Leu	Glu 670	Glu	Ile

Trp	Asp	Val 675	His	Ala	Ile	Pro	Pro 680	Ser	Val	Gln	Ser	His 685	Ile	Gln	Ala	
Thr	Gln 690	Ala	Glu	Lys	Ala	Ala 695	Gln	Glu	Ala	Gln	Ala 700	Thr	Thr	Ser	Ala	
Ile 705	Ser	Ala	Ala	Ala	Thr 710	Ser	Ser	Ser	Ser	Ile 715	Asn	Thr	Ser	Met	Ala 720	
Thr	Ser	Ser	Ser	Ser 725	Ser	Leu	Ser	Pro	Ser 730	Ala	Ala	Ser	Thr	Pro 735	Asn	
Gly	Gly	Ala	Val 740	Asp	Tyr	Val	GJ.y	Thr 745	Asp	Met	Ser	Met	Ser 750	Leu	Val	
Gln	Ser	Asp 755	Asn	Ala												
<213 <213	0> 3 l> 14 2> DN 3> Lu	JA.	ia cı	ıprin	na									,		
)> L> CI 2> (1		(140)	L)												
<400) ~ 2															
atg	gat Asp									_	_	_	_	_		48
	cag Gln			Lys	Pro	Asp		Ser	Leu			_			_	96
	agt Ser															144
	cag Gln 50															192
	atg Met	_			_				_	_	_					240

	ctt Leu														288
	gga Gly														336
	aaa Lys 115														384
	aat Asn														432
	tat Tyr		Lys												480
	gaa Glu	_		_			_	_	_		_	_	_	_	528
	ggc Gly													-	576
	ggc Gly 195								-			-	_		624
	gaa Glu								_	-	-		_	-	672
	aac Asn														720
	gac Asp					_				_	_	_	_		768
	ctc Leu			_	_	-		_	_	_					816
	ttg Leu 275	_	-		_	_			_		_	_			864

aat Asn	gaa Glu 290	ctg Leu	cta Leu	att Ile	gca Ala	aat Asn 295	gtt Val	gcc Ala	tgg Trp	tgc Cys	agt Ser 300	att Ile	gag Glu	tct Ser	ctg Leu	912
		gaa Glu														960
		tca Ser														1008
		tat Tyr														1056
		cgt Arg 355			Ser		_	_			_		-			1104
		cgc Arg														1152
		ata Ile														1200
		atc Ile											_			1248
		gat Asp														1296
		atc Ile 435														1344
		gaa Glu											-	_	_	1392
	atc Ile	_														1401

<21 <21	0> 4 1> 4 2> P: 3> L:	RT	ia c	upri	na										
-40	0 > 4														
		Asn	Gly	Glu 5	Gln	Asp	Ala	Gly	Phe 10	Arg	Leu	Ala	Pro	Met 15	Ser
Pro	Gln	Glu	Ile 20	Lys	Pro	Asp	Ile	Ser 25	Leu	Leu	Asn	Glu	Asn 30	Asn	Thr
Ser	Ser	Tyr 35	Ser	Pro	Lys	Pro	Gly 40	Ser	Pro	Asn	Pro	Phe 45	Ala	Ile	Gly
Leu	Gln 50	Ala	Ile	Asn	Ala	Val 55	Ala	Ala	Ala	Asn	Ala 60	Asn	Asn	Gln	Asn
Gln 65	Met	Leu	Gln	Thr	Thr 70	Pro	Pro	Gln	Gln	Gln 75	Gln	Tyr	Pro	Pro	Asn 80
His	Pro	Leu	Ser	Gly 85	Ser	Lys	His	Leu	Cys 90	Ser	Ile	Cys	Gly	Asp 95	Arg
Ala	Ser	Gly	Lys 100	His	Tyr	Gly	Val	Tyr 105	Ser	Cys	Glu	Gly	Cys 110	Lys	Gly
Phe	Phe	Lys 115	Arg	Thr	Val	Arg	Lys 120	Asp	Leu	Thr	Tyr	Ala 125	Cys	Arg	Glu
Asp	Arg 130	Asn	Cys	Ile	Ile	Asp 135	Lys	Arg	Gln	Arg	Asn 140	Arg	Cys	Gln	Tyr
Cys 145	Arg	Tyr	Gln	Lys	Cys 150	Leu	Ala	Cys	Gly	Met 155	Lys	Arg	Glu	Ala	Val 160
Gln	Glu	Glu	Arg	Gln 165	Arg	Gly	Thr	Arg	Ala 170	Ala	Asn	Ala	Arg	Ala 175	Ala
Gly	Ala	Gly	Gly 180	Gly	Gly	Gly	Gly	Gly 185	Gly	Gly	Val	Ser	Asn 190	Val	Val
Gly	Ala	Gly 195	Gly	Glu	Asp	Phe	Lys 200	Pro	Ser	Ser	Ser	Leu 205	Arg	Asp	Leu
Thr	Ile 210	Glu	Arg	Ile	Ile	Glu 215	Ala	Glu	Gln	Lys	Ala 220	Glu	Ser	Leu	Ser
Gly 225	Asp	Asn	Val	Leu	Pro 230	Phe	Leu	Arg	Val	Gly 235	Asn	Asn	Ser	Met	Val 240

Gln His Asp Tyr Lys Gly Ala Val Ser His Leu Cys Gln Met Val Asn 245 250 Lys Gln Leu Tyr Gln Met Val Glu Tyr Ala Arg Arg Thr Pro His Phe 260 265 Thr His Leu Gln Arg Glu Asp Gln Ile Leu Leu Leu Lys Ala Gly Trp 280 Asn Glu Leu Leu Ile Ala Asn Val Ala Trp Cys Ser Ile Glu Ser Leu 290 295 Asp Ala Glu Tyr Ala Ser Pro Gly Thr Val His Asp Gly Ser Phe Gly 305 310 Arg Arg Ser Pro Val Arg Gln Pro Gln Gln Leu Phe Leu Asn Gln Asn Phe Ser Tyr His Arg Asn Ser Ala Ile Lys Ala Asn Val Val Ser Ile 345 Phe Asp Arg Ile Leu Ser Glu Leu Ser Ile Lys Met Lys Arg Leu Asn 355 360 Ile Asp Arg Ser Glu Leu Ser Cys Leu Lys Ala Ile Ile Leu Phe Asn 370 375 Pro Asp Ile Arg Gly Leu Lys Cys Arg Ala Asp Val Glu Val Cys Arg 390 Glu Lys Ile Tyr Ala Cys Leu Asp Glu His Cys Arg Thr Glu His Pro 405 410 Gly Asp Asp Gly Arg Phe Ala Gln Leu Leu Leu Arg Leu Pro Ala Leu 420 425 Arg Ser Ile Ser Leu Lys Cys Leu Asp His Leu Phe Phe Phe Arg Leu 435 440 Ile Gly Glu Arg Ala Leu Glu Glu Leu Ile Ala Glu Gln Leu Glu Ala 450 455 460 Pro Ile Cys 465 <210> 5 <211> 585 <212> DNA <213> Myzus persicae

	1> C	DS 1)	(585)										
gaa		ggc Gly												48
		ctt Leu												96
		ttg Leu 35							-		_	_	_	144
		ggc								_			_	192
		tgc Cys								_		-		240
		tgc Cys				_	_		_	-		_		288
		tgc Cys												336
		gaa Glu 115												384
		aaa Lys												432
		cct Pro								-	_		_	480
_		gaa Glu		_	_			_	_		_			528

gtg aaa cct Val Lys Pro tac aaa ttt Tyr Lys Phe 195											
<210> 6 <211> 195 <212> PRT <213> Myzus	persica	e									
<400> 6 Glu Phe Gly 1	Thr Ser	Ala Il	e Val	Asn	Gly 10	Phe	Ile	Arg	Thr	Ile 15	Ser
Leu Ile Leu	Ile Phe 20	Leu Le	ı Leu	Phe 25	Leu	Trp	Arg	Leu	Leu 30	Ala	Phe
Arg Phe Leu 35	Phe Ile	Ser Gl	ı Gln 40	Pro	Pro	Pro	Glu	Glu 45	Leu	Cys	Leu
Val Cys Gly 50	Asp Arg	Ser Ser		Tyr	His	Tyr	Asn 60	Ala	Leu	Thr	Cys
Glu Gly Cys 65	Lys Gly	Phe Phe	e Arg	Arg	Ser	Ile 75	Thr	Lys	Asn	Ala	Val 80
Tyr Gln Cys	Lys Tyr 85	Gly Ası	ı Asn	Cys	Glu 90	Ile	Asp	Met	Tyr	Met 95	Arg
Arg Lys Cys	Gln Glu 100	Cys Arg	g Leu	Lys 105	Lys	Cys	Leu	Thr	Val 110	Gly	Met
Arg Pro Glu 115	Cys Val	Val Pro	Glu 120	Val	Gln	Cys	Ala	Val 125	Lys	Arg	Lys
Glu Lys Lys 130	Ala Gln	Arg Glu		Asp	Lys	Pro	Asn 140	Ser	Thr	Thr	Asp
Ile Ser Pro 145	Glu Ile	Ile Lys 150	: Ile	Glu	Pro	Thr 155	Glu	Met	Lys	Ile	Glu 160
Cys Gly Glu	Pro Met 165	Ile Met	: Gly	Thr	Pro 170	Met	Pro	Thr	Val	Pro 175	Tyr
Val Lys Pro	Leu Ser 180	Ser Le	ı Val	Pro 185	Asn	Ser	Ala	Arg	Val 190	Thr	Gly

Tyr Lys Phe 195

<210> 7 <211> 208 <212> DNA

<213> Myzus persicae

<400> 7
catgcctgca ggtcgactct agaggatccc ctcgtccggt taccattaca acgcactcac 60
ctgtgaaggc tgtaagggtt tctttcgacg gagtgttacc aaaaatgcgg tgtattgttg 120
taaatttggt catgcctgcg aaatggacat gtatatgcga cgtaaatgtc aggaatgtag 180
gctgaaaaaa tgtttggctg tgggcatg
208

<210> 8 <211> 436 <212> DNA <213> Myzus persicae

<400> 8
catgcggccg gaatgtgtgg tgcccgaaaa ccagtgtgca atgaaacgac gcgaaaagaa 60
agcacaaaaa gagaaggata aaatacagac cagtgtgtg gcaacggaaa ttaaaaagga 120
aatactcgat ttaatgacat gtgaaccgcc atcacatcca acgtgtccgc tgttacctga 180
agacattttg gctaaatgtc aagctcgtaa tatacctcct ttatcgtaca atcaattggc 240
agttatatat aaattaatat ggtatcaaga tggctacgaa cagccatccg aggaagatct 300
caaacgtata atgagttcac ccgatgaaaa tgaaagtcaa cacgatgcat catttcgtca 360
tataacagaa atcactatac taacagtaca attaattgtt gaatgtgcca aaggtctagg 420
gtaccgagct cgaatt

<210> 9
<211> 1797
<212> DNA
<213> Myzus persicae
<220>
<221> CDS
<222> (1)..(1797)

	0> 9															
atg Met 1	atg Met	gac Asp	cag Gln	aaa Lys 5	tgt Cys	gac Asp	gtc Val	ggc	ggt Gly 10	ggt	ggt Gly	gtc Val	gct Ala	gct Ala 15	gcc Ala	48
gcc Ala	gcc Ala	ggt Gly	atc Ile 20	ggt Gly	ggc Gly	ggc Gly	ggt Gly	gtc Val 25	ggc Gly	ggc	ctc Leu	atg Met	tcg Ser 30	tac Tyr	aac Asn	96
cgt Arg	ggc	cgt Arg 35	ggc	ggc Gly	acc Thr	gag Glu	gtc Val 40	atc Ile	atc Ile	aaa Lys	ccc Pro	cgt Arg 45	agt Ser	cct Pro	gcc Ala	144
gtg Val	gtg Val 50	cag Gln	gtg Val	gcc Ala	acc Thr	ggt Gly 55	ggc Gly	agt Ser	tac Tyr	cac His	ggc Gly 60	ctg Leu	ccg Pro	gcg Ala	gcc Ala	192
					gtg Val 70											240
					ccg Pro											288
					gtc Val											336
gtc Val	aac Asn	tcg Ser 115	ccg Pro	ccc Pro	gcc Ala	tct Ser	tcg Ser 120	ccc Pro	ggc	acg Thr	tcg Ser	cac His 125	ata Ile	tcc Ser	tac Tyr	384
			Ser	Asn	ggc Gly	Gly										432
					atg Met 150											480
					gtc Val											528
					cat His											576

cac His	acg Thr	ccc Pro 195	tcg Ser	acc Thr	ggt Gly	gtc Val	gtc Val 200	aac Asn	acc Thr	tcg Ser	gca Ala	tcg Ser 205	ggc Gly	ccc Pro	Gly 999	624
ggt Gly	ggc Gly 210	gtt Val	ggt Gly	ggc	aat Asn	gtg Val 215	ctg Leu	aac Asn	aac Asn	cga Arg	cct Pro 220	ccc Pro	gaa Glu	gag Glu	ctg Leu	672
tgc Cys 225	ctg Leu	gtg Val	tgt Cys	ggc	gac Asp 230	cgg Arg	tcg Ser	tcc Ser	ggt Gly	tac Tyr 235	cat His	tac Tyr	aac Asn	gct Ala	ctc Leu 240	720
					aag Lys											768
				Cys	aag Lys											816
					cag Gln											864
					tgt Cys											912
					gct Ala 310											960
					gaa Glu											1008
att Ile	gaa Glu	tgt Cys	ggt Gly 340	gaa Glu	cca Pro	atg Met	ata Ile	atg Met 345	Gly	aca Thr	cct Pro	atg Met	ccg Pro 350	act Thr	gta Val	1056
					ttg Leu											1104
					gat Asp											1152
					aat Asn 390				-	_	_		_	_	_	1200

aaa Lys	caa Gln	agt Ser	gac Asp	acc Thr 405	aca Thr	tat Tyr	cga Arg	atc Ile	atc Ile 410	act Thr	gag Glu	atg Met	aca Thr	ata Ile 415	ctc Leu	1248
					gtt Val											1296
					gat Asp											1344
					agg Arg											1392
				Ala	aac Asn 470											1440
					gat Asp											1488
					aag Lys											1536
					tcg Ser											1584
					gaa Glu				_				_		_	1632
					gac Asp 550											1680
					ttg Leu											1728
					ctg Leu					-				_	_	1776
_			_	_	atg Met											1797

<210> 10 <211> 599						
<211> 399						
<213> Myzus	nergicae					
(213) MyZus	persicae					
<400> 10						
Met Met Asp	Gln Lys C	ys Asp Va	l Gly Gly	Gly Gly	Val Ala	Ala Ala
1	5		10			15
Ala Ala Gly		TA GIA GI		Gly Leu		Tyr Asn
	20		25		30	
Arg Gly Arg	Gly Gly T	hr Glu Va	l Tla Tla	Tug Dro	Ara Cor	Dwo Nlo
35	GIY GIY I	4:		nys FIO	45	PIO AIA
33		•	•		4.0	
Val Val Gln	Val Ala T	hr Gly Gl	Ser Tyr	His Gly	Leu Pro	Ala Ala
50		55	•	60		
Ser Asp Ala	Val Ile V	al Arg Se	r Pro Pro	Gly Gly	His Leu	Pro Gly
65		70		75		80
Pro Gln Gln		ro Pro Se	_	Gly Cys	Ser Thr	
	85		90			95
C 7 T1-	71 - Gl 11	. 7 * 7		D 3		
Ser Asp Ile	=	at Lys Arg	-	Pro Asp		Leu Ala
	100		105		110	
Val Asn Ser	Pro Pro A	la Ser Sei	r Pro Glv	Thr Ser	His Tle	Ser Tur
115	110 110 A	120	_	1111 501	125	Der Tyt
Thr Val Ile	Ser Asn G	ly Gly Gly	Gly Gly	Gly Gly	Gly Gly	Gly Gly
130		135		140		
Tyr Asn Thr	Ser Pro M	et Ser Thi	Asn Ser	Tyr Asp	Pro Tyr	Ser Pro
145	1.	50		155		160
					_	
Met Ser Gly		al Lys Gli		Ser Pro	Pro Asn	
	165		170			175
Ser Gly Val	Ser Ser W	ie Ser Ner	Cly Leu	Lve Lve	Lve Lve	Leu Asn
ber dry var	180	ra der wal	185	nya nya	190	Dea Asii
	100		203		200	
His Thr Pro	Ser Thr G	ly Val Val	Asn Thr	Ser Ala	Ser Gly	Pro Gly
195		200			205	•
Gly Gly Val	Gly Gly A	sn Val Le	ı Asn Asn	Arg Pro	Pro Glu	Glu Leu
210		215		220		
					 -	
Cys Leu Val		-	Ser Gly		Tyr Asn	
225	2	30		235		240

Thr	Cys	Glu	Gly	Cys 245	Lys	Gly	Phe	Phe	Arg 250	Arg	Ser	Ile	Thr	Lys 255	Asn
Ala	Val	Tyr	Gln 260	Cys	ГЛЗ	Tyr	Gly	Asn 265	Asn	Cys	Glu	Ile	Asp 270	Met	Tyr
Met	Arg	Arg 275	Lys	Cys	Gln	Glu	Cys 280	Arg	Leu	Lys	Lys	Cys 285	Leu	Thr	Val
Gly	Met 290	Arg	Pro	Glu	Cys	Val 295	Val	Pro	Glu	Val	Gln 300	Cys	Ala	Val	Lys
Arg 305	Lys	Glu	Lys	Lys	Ala 310	Gln	Arg	Glu	Lys	Asp 315	Lys	Pro	Asn	Ser	Thr 320
Thr	Asp	Ile	Ser	Pro 325	Glu	Ile	Ile	Lys	Ile 330	Glu	Pro	Thr	Glu	Met 335	Lys
Ile	Glu	Cys	Gly 340	Glu	Pro	Met	Ile	Met 345	Gly	Thr	Pro	Met	Pro 350	Thr	Val
Pro	Tyr	Val 355	Lys	Pro	Leu	Ser	Ser 360	Glu	Gln	Lys	Glu	Leu 365	Ile	His	Arg
Leu	Val 370	Tyr	Phe	Gln	Asp	Gln 375	Tyr	Glu	Ala	Pro	Ser 380	Glu	Lys	Asp	Met
Lys 385	Arg	Leu	Thr	Ile	Asn 390	Asn	Gln	Asn	Met	Asp 395	Glu	Tyr	Asp	Glu	Glu 400
Lys	Gln	Ser	Asp	Thr 405	Thr	Tyr	Arg	Ile	Ile 410	Thr	Glu	Met	Thr	Ile 415	Leu
Thr	Val	Gln	Leu 420	Ile	Val	Glu	Phe	Ala 425	Lys	Arg	Leu	Pro	Gly 430	Phe	Asp
Lys	Leu	Val 435	Arg	Glu	Asp	Gln	Ile 440	Thr	Leu	Leu	Lys	Ala 445	Cys	Ser	Ser
Glu	Ala 450	Met	Met	Phe	Arg	Val 455	Ala	Arg	Lys	Tyr	Asp 460	Ile	Thr	Thr	Asp
Ser 465	Ile	Val	Phe	Ala	Asn 470	Asn	Gln	Pro	Phe	Ser 475	Ala	Asp	Ser	Tyr	Asn 480
Lys	Ala	Gly	Leu	Gly 485	Asp	Ala	Ile	Glu	Asn 490	Gln	Leu	Ser	Phe	Ser 495	Arg
Phe	Met	Tyr	Asn 500	Met	Lys	Val	Asp	Asn 505	Ala	Glu	Tyr	Ala	Leu 510	Leu	Thr

515 520	Leu Asp Gly Trp Lys 525
Val Glu Lys Ile Gln Glu Ile Tyr Leu Glu Ser 530 535	Leu Lys Ala Tyr Val 540
Asp Asn Arg Asp Arg Asp Thr Ala Thr Val Arg	-
Ser Val Leu Thr Glu Leu Arg Thr Leu Gly Asn 565 570	Glu Asn Ser Glu Leu 575
Cys Met Thr Leu Lys Leu Lys Asn Arg Val Val 580 585	Pro Pro Phe Leu Ala 590
Glu Ile Trp Asp Val Met Pro 595	
<210> 11 <211> 1131 <212> DNA <213> Myzus persicae	
<220> <221> CDS <222> (1)(1131)	
<pre><400> 11 atg tat tcc aac tcg tac acc atg tat tca agt Met Tyr Ser Asn Ser Tyr Thr Met Tyr Ser Ser</pre>	
atg tat tcc aac tcg tac acc atg tat tca agt Met Tyr Ser Asn Ser Tyr Thr Met Tyr Ser Ser	Asp Arg Leu Tyr Ser 15 aac gta caa gac tct 96
atg tat tcc aac tcg tac acc atg tat tca agt Met Tyr Ser Asn Ser Tyr Thr Met Tyr Ser Ser 1 5 5 10 gtc gat cgg aac agt atg atg aat aat tct tgc Val Asp Arg Asn Ser Met Met Asn Asn Ser Cys	Asp Arg Leu Tyr Ser 15 aac gta caa gac tct 96 Asn Val Gln Asp Ser 30 tcg aaa cat ctg tgc 144
atg tat tcc aac tcg tac acc atg tat tca agt Met Tyr Ser Asn Ser Tyr Thr Met Tyr Ser Ser 1 5 5 10 gtc gat cgg aac agt atg atg aat aat tct tgc Val Asp Arg Asn Ser Met Met Asn Asn Ser Cys 20 25 ccg aat tac ccg ccc aac cat cca ctc agc ggt Pro Asn Tyr Pro Pro Asn His Pro Leu Ser Gly	Asp Arg Leu Tyr Ser 15 aac gta caa gac tct 96 Asn Val Gln Asp Ser 30 tcg aaa cat ctg tgc 144 Ser Lys His Leu Cys 45 tac gga gtc tac agc 192
atg tat tcc aac tcg tac acc atg tat tca agt Met Tyr Ser Asn Ser Tyr Thr Met Tyr Ser Ser 1	Asp Arg Leu Tyr Ser 15 aac gta caa gac tct 96 Asn Val Gln Asp Ser 30 tcg aaa cat ctg tgc 144 Ser Lys His Leu Cys 45 tac gga gtc tac agc 192 Tyr Gly Val Tyr Ser 60 gtg agg aaa aat ttg 240

		cgg Arg														336
		aga Arg 115														384
		aat Asn														432
		gtg Val	_					_				_	_	_		480
		gaa Glu		-	•		_		-						_	528
		att Ile														576
		aag Lys 195														624
		tta Leu								_	_		-	_		672
		aga Arg						_			-		_			720
		gtt Val										_	_	-		768
	-	cgt Arg	_			-		_	_		_	_	_	-		816
_	_	aga Arg 275						-	_							864
cca	ggt	tca	aaa	ggt	ttg	cag	tct	gtg	aat	gaa	gtg	caa	gta	ctg	cgt	912

Pro Gly Ser Lys Gly Leu Gln Ser Val Asn Glu Val Gln Val Leu Arg 290 295 300	
gat aag gtt tat gtt gcg tta gaa gaa tat tgt cgt aca aca cat cca Asp Lys Val Tyr Val Ala Leu Glu Glu Tyr Cys Arg Thr Thr His Pro 305 310 315 320	960
gaa gaa cct gga cga ttt gct aaa cta ctt ctt cgg ctt cct tca tta Glu Glu Pro Gly Arg Phe Ala Lys Leu Leu Leu Arg Leu Pro Ser Leu 325 330 335	1008
cgt tca att gga tta aaa tgt ctg gaa cat tta ttc ttt tat aaa ctt Arg Ser Ile Gly Leu Lys Cys Leu Glu His Leu Phe Phe Tyr Lys Leu 340 345 350	1056
att ggc gat tcc cca att gat aca ttt tta atg gaa gtt ctc gaa tca Ile Gly Asp Ser Pro Ile Asp Thr Phe Leu Met Glu Val Leu Glu Ser 355 360 365	1104
tct tca cat gac gtt caa gta gct aca Ser Ser His Asp Val Gln Val Ala Thr 370 375	· 1131
<210> 12	
<211> 377 <212> PRT <213> Myzus persicae	
<212> PRT	
<212> PRT <213> Myzus persicae <400> 12 Met Tyr Ser Asn Ser Tyr Thr Met Tyr Ser Ser Asp Arg Leu Tyr Ser	
<pre><212> PRT <213> Myzus persicae <400> 12 Met Tyr Ser Asn Ser Tyr Thr Met Tyr Ser Ser Asp Arg Leu Tyr Ser</pre>	
<pre><212> PRT <213> Myzus persicae <400> 12 Met Tyr Ser Asn Ser Tyr Thr Met Tyr Ser Ser Asp Arg Leu Tyr Ser</pre>	
<pre><212> PRT <213> Myzus persicae <400> 12 Met Tyr Ser Asn Ser Tyr Thr Met Tyr Ser Ser Asp Arg Leu Tyr Ser 1</pre>	
<pre><212> PRT <213> Myzus persicae <400> 12 Met Tyr Ser Asn Ser Tyr Thr Met Tyr Ser Ser Asp Arg Leu Tyr Ser 1</pre>	

Met	Lys	Arg 115		Ala	Val	Gln	Glu 120	Glu	Arg	Gln	Arg	Thr 125	Lys	Glu	Arg
Asp	His 130	Asn	Asn	Ile	Glu	Val 135	Glu	Pro	Thr	Ser	Ser 140	Ser	Asn	Thr	Asp
Met 145	Pro	Val	Glu	Leu	Ile 150	Leu	Arg	Ala	Glu	Asn 155	Lys	Ala	Asp	Ala	Ile 160
Lys	Thr	Glu	Gln	Gln 165	Tyr	Ile	Glu	Gln	Arg 170	His	Pro	Gln	His	Thr 175	Val
Gly	Ala	Ile	Cys 180	Gln	Ala	Thr	Asp	Lys 185	Gln	Leu	Ile	Gln	Leu 190	Val	Glu
Trp	Ala	Lys 195	His	Ile	Pro	His	Phe 200	Lys	Asn	Leu	Pro	Leu 205	Gly	Asp	Gln
Val	Leu 210	Leu	Leu	Arg	Ala	Gly 215	Trp	Asn	Glu	Leu	Met 220	Ile	Ala	Ala	Phe
Ser 225	His	Arg	Ser	Ile	Ser 230	Val	Lys	Asp	Gly	Ile 235	Val	Leu	Ala	Thr	Gly 240
Leu	Thr	Val	Asp	Arg 245	Asp	Ser	Ala	His	Gln 250	Ala	Gly	Val	Glu	Ala 255	Ile
Phe	Asp	Arg	Val 260	Leu	Thr	Glu	Leu	Val 265	Ala	Lys	Met	Arg	Asp 270	Met	Gly
Met	Asp	Arg 275	Thr	Glu	Leu	Gly	Cys 280	Leu	Arg	Thr	Ile	Ile 285	Leu	Phe	Asn
Pro	Gly 290	Ser	Lys	Gly	Leu	Gln 295	Ser	Val	Asn	Glu	Val 300	Gln	Val	Leu	Arg
Asp 305	Lys	Val	Tyr	Val	Ala 310	Leu	Glu	Glu	Tyr	Cys 315	Arg	Thr	Thr	His	Pro 320
Glu	Glu	Pro	Gly	Arg 325	Phe	Ala	Lys	Leu	Leu 330	Leu	Arg	Leu	Pro	Ser 335	Leu
Arg	Ser	Ile	Gly 340	Leu	Lys	Cys	Leu	Glu 345	His	Leu	Phe	Phe	Tyr 350	Lys	Leu
Ile	Gly	Asp 355	Ser	Pro	Ile	Asp	Thr 360	Phe	Leu	Met	Glu	Val 365	Leu	Glu	Ser
Ser	Ser 370	His	Asp	Val	Gln	Val 375	Ala	Thr							

```
<210> 13
<211> 150
<212> DNA
<213> Lucilia cuprina
<220>
<221> CDS
<222> (9)..(134)
<400> 13
aattetge gaa gga tge aag gga tte tte aaa egt ace gta ege aag gae 50
         Glu Gly Cys Lys Gly Phe Phe Lys Arg Thr Val Arg Lys Asp
ttg aca tat gct tgt cgt gag gac aga aat tgc att ata gat aaa cga
                                                                   98
Leu Thr Tyr Ala Cys Arg Glu Asp Arg Asn Cys Ile Ile Asp Lys Arg
 15
                     20
                                          25
caa aga aat cgt tgc cag tat tgt cgc tac caa aag tgatcgatac cqtcqa 150
Gln Arg Asn Arg Cys Gln Tyr Cys Arg Tyr Gln Lys
                 35
<210> 14
<211> 42
<212> PRT
<213> Lucilia cuprina
<400> 14
Glu Gly Cys Lys Gly Phe Phe Lys Arg Thr Val Arg Lys Asp Leu Thr
Tyr Ala Cys Arg Glu Asp Arg Asn Cys Ile Ile Asp Lys Arg Gln Arg
Asn Arg Cys Gln Tyr Cys Arg Tyr Gln Lys
         35
<210> 15
<211> 32
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence:oligonucleotide
      useful as primer.
<220>
<221> misc_feature
<222> (1)..(32)
```

<223> Nucleotides designated as "n" residues can be A, G, C or T residues.	
<400> 15 cggaattccg cctcnggnta ycaytayaay gc	32
<210> 16 <211> 32 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence:Oligonucleotide useful as primer.	
<400> 16	
cgcggatccr cactcctgac actttcgyct ca	32
<210> 17	-
<211> 23	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:Oligonucleotide useful as primer.	
<400> 17	
gcctcggggt atcactataa cgc	23
<210> 18	
<211> 23	
<212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence:Oligonucleotide useful as primer.	
<400> 18	
gcactcctga cactttcgtc tca	23
<210× 10	
<210> 19 <211> 23	
<212> DNA	
<213> Artificial Sequence	

<220> <223> Description of Artificial Sequence:Oligonucleotide useful as primer. <400> 19 tcgtccggtt accattacaa cgc 23 <210> 20 <211> 24 <212> DNA <213> Artificial Sequence <220> <221> misc_feature <222> (1)..(24) <223> Nucleotides designated as "n" residues can be A, G, C or T residues. <400> 20 tagacctttg gcraaytcna caat 24